AMENDMENTS TO THE CLAIMS

1-4. (cancelled)

- 5. (currently amended) A plasmid for expression of recombinant eukaryotic genes, the plasmid comprising:
 - a first transcription unit comprising a first transcriptional control sequence transcriptionally linked with a first 5'-untranslated region comprising a first synthetic intron, a first coding sequence, and a first 3'-untranslated region/poly(A) signal, wherein said first synthetic intron is between said first transcriptional control sequence and said first coding sequence; and
 - a second transcription unit comprising a second transcriptional control sequence transcriptionally linked with a second 5'-untranslated region comprising a second synthetic intron, a second coding sequence, and a second 3'-untranslated region/poly(A) signal, wherein said second synthetic intron is between said second transcriptional control sequence and said second coding sequence,
 - wherein the first and second synthetic introns both comprise a 5' splice site having [[a]] the sequence of residues 1-9 of SEQ ID NO:13, a branch point having [[a]] the sequence of residues 93-99 of SEQ ID NO:13, and a 3' splice site having [[a]] the sequence of residues 102-122 of SEQ ID NO:13.

6-9. (cancelled)

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- 10. (currently amended) A plasmid for expression of recombinant enkaryotic genes, the plasmid comprises:
 - a transcriptional control sequence transcriptionally linked with a first coding sequence and a second coding sequence;
 - a 5'-untranslated region;
 - an intron 5' to said first coding sequence, wherein the intron comprises a 5' splice site having [[a]] the sequence of residues 1-9 of SEQ ID NO:13, a branch point having [[a]] the sequence of residues 93-99 of SEQ ID NO:13, and a 3' splice site having [[a]] the sequence of residues 102-122 of SEQ ID NO:13;
 - an alternative 3' splice site located between the first and second coding sequence; and a 3'-untranslated region/poly(A) signal.

11-13. (cancelled)

- 14. (currently amended) A plasmid for expression of recombinant eukaryotic genes comprising:
 - a transcriptional control sequence transcriptionally linked with a first coding sequence, an IRES sequence, a second coding sequence, and a 3'-untranslated region/poly(A) signal, wherein said IRES sequence is between said first coding sequence and said second coding sequence; and
 - a synthetic intron between said transcriptional control sequence and said first coding sequence, wherein the synthetic intron comprises a 5' splice site having [[a]] the sequence of residues 1-9 of SEQ ID NO:13, a branch point having [[a]] the sequence of residues 93-99 of SEQ ID NO:13, and a 3' splice site having [[a]] the sequence of residues 102-122 of SEQ ID NO:13.

15-50. (cancelled)

51. (currently amended) The plasmid of claim 10 wherein both the 3' splice site and the alternative 3' splice site have [[a]] the sequence of residues 102-122 of SEQ ID NO:13.

- 52. (previously presented) The plasmid of claim 10 wherein the 3' splice site is weakened with respect to the alternative 3' splice site.
- 53. (previously presented) The plasmid of claim 10 wherein the 3' splice site is weakened with respect to the alternative 3' splice site by changing three consecutive T's to A's.
- 54. (currently amended) The plasmid of claim 10 wherein the 3' splice site has [[a]] the sequence of residues 102-122 of SEQ ID NO:13 with residues 108-110 replaced by AAA and the alternative 3' splice site has [[a]] the sequence of residues 102-122 of SEQ ID NO:13.

55-64. (canceled)

- 65. (currently amended) A synthetic transcription unit for efficient and accurate expression of recombinant eukaryotic genes, the transcription unit comprising a synthetic intron comprising:
 - a 5' splice site having [[a]] the sequence of SEQ ID NO:15, a branch point having [[a]] the sequence of SEQ ID NO:17, and a 3' splice site having [[a]] the sequence of Y₁₆NYAGG, wherein Y = C or T, Y₁₆ contains 7 consecutive T residues, and N = any base. SEQ-ID NO:18, wherein the 3' splice site contains 7 consecutive T residues.

66-68. (canceled)

69. (currently amended) The synthetic transcription unit of claim 65, wherein the branch point and the 3' splice site together have [[a]] the sequence of TACTAACGGTTCTTTTTTCTCTTCACAGG (SEQ ID NO:13, residues #93 through #122).

70-71. (canceled)

72. (previously presented) A synthetic transcription unit for efficient and accurate expression of recombinant eukaryotic genes, wherein the transcriptional unit comprises a synthetic intron having the sequence of SEQ ID NO:13.

73-76. (canceled)